

6 TRANSOUENCE LISTING															
<110>	Fujis Nakaj Nishi Nishi Yoshi Ohkub	ima, o, K o, N mura	Hid aori Iobuy 1, Se	lenor va eiji		al C	lo.,	Ltd							
<120>	Novel	35	kD F	rote	in										
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_	_	Phe		_	aac Asn	_				gggc	agg	cagg	gtcc	at		1008
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1 10 15

Ser Leu Ser Arg Asn Val Gly Val Trp Ala Ser Gly Glu Gly Lys Lys 20 25 30

Val Asp Ile Ala Gly Ile Tyr Pro Pro Val Thr Thr Pro Phe Thr Ala 35 40 45

Thr Ala Glu Val Asp Tyr Gly Lys Leu Glu Glu Asn Leu His Lys Leu 50 60

Gly Thr Phe Pro Phe Arg Gly Phe Val Val Gln Gly Ser Asn Gly Glu 65 70 75 80

Phe Pro Phe Leu Thr Ser Ser Glu Arg Leu Glu Val Val Ser Arg Val
85 90 95

Arg Gln Ala Met Pro Lys Asn Arg Leu Leu Leu Ala Gly Ser Gly Cys
100 105 110

Glu Ser Thr Gln Ala Thr Val Glu Met Thr Val Ser Met Ala Gln Val 115 120 125

Gly Ala Asp Ala Ala Met Val Val Thr Pro Cys Tyr Tyr Arg Gly Arg 130 135 140

Met Ser Ser Ala Ala Leu Ile His His Tyr Thr Lys Val Ala Asp Leu 145 150 155 160

Ser Pro Ile Pro Val Val Leu Tyr Ser Val Pro Ala Asn Thr Gly Leu 165 170 175

Asp Leu Pro Val Asp Ala Val Val Thr Leu Ser Gln His Pro Asn Ile 180 185 190

Val Gly Met Lys Asp Ser Gly Gly Asp Val Thr Arg Ile Gly Leu Ile 195 200 205 Val His Lys Thr Arg Lys Gln Asp Phe Gln Val Leu Ala Gly Ser Ala Gly Phe Leu Met Ala Ser Tyr Ala Leu Gly Ala Val Gly Gly Val Cys Ala Leu Ala Asn Val Leu Gly Ala Gln Val Cys Gln Leu Glu Arg Leu Cys Cys Thr Gly Gln Trp Glu Asp Ala Gln Lys Leu Gln His Arg Leu Ile Glu Pro Asn Ala Ala Val Thr Arg Arg Phe Gly Ile Pro Gly Leu Lys Lys Ile Met Asp Trp Phe Gly Tyr Tyr Gly Gly Pro Cys Arg Ala Pro Leu Gln Glu Leu Ser Pro Ala Glu Glu Glu Ala Leu Arg Met Asp Phe Thr Ser Asn Gly Trp Leu <210> <211> <212> DNA <213> Rattus sp. <220> <221> CDS <222> (9)..(1016) <400> 3 cgggatcc atg ctg ggc ccc caa atc tgg gcc tcc atg agg cag ggg ctg Met Leu Gly Pro Gln Ile Trp Ala Ser Met Arg Gln Gly Leu agc agg ggc ttg tct agg aac gtg aag ggg aag aag ata gac att gcc Ser Arg Gly Leu Ser Arg Asn Val Lys Gly Lys Lys Ile Asp Ile Ala ggc atc tac cca ccc gtg acc acc cca ttc acc gcc acc gca gaa gta Gly Ile Tyr Pro Pro Val Thr Thr Pro Phe Thr Ala Thr Ala Glu Val gac tat ggg aaa ctg gaa gag aac ctg aac aaa ctg gcc gcc ttc ccc Asp Tyr Gly Lys Leu Glu Glu Asn Leu Asn Lys Leu Ala Ala Phe Pro ttt cga ggc ttc gtg gtc cag ggc tct act gga gag ttt cca ttc ctg Phe Arg Gly Phe Val Val Gln Gly Ser Thr Gly Glu Phe Pro Phe Leu

65						75					
acc Thr	•		-								

	.—			_				gtg Val	_	<del>-</del>	-				290
	_	_			_		_	ggc Gly			_		_		338
_		_		_		_		atg Met	_		· <del>-</del>				386
_	_					_		tat Tyr 135	_		_			_	434
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Val								aac Asn		_				_	530
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								act Thr							626
								gct Ala 215						ctg Leu	674
								ggg Gly							722
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· ·								cag Gln							818
	_							ata Ile							866
_								ccc Pro 295							914
			Ser									_		aat Asn	962

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Val Val Thr Leu Ser Gln His Pro Asn Ile Ile Gly Leu Lys Asp Ser 180 185 190

Gly Gly Asp Val Thr Arg Thr Gly Leu Ile Val His Lys Thr Ser Lys
195 200 205

Gln Asp Phe Gln Val Leu Ala Gly Ser Val Gly Phe Leu Leu Ala Ser 210 220

Tyr Ala Val Gly Ala Val Gly Gly Ile Cys Gly Leu Ala Asn Val Leu 225 230 235 240

Gly Ala Gln Val Cys Gln Leu Glu Arg Leu Cys Leu Thr Gly Gln Gly
245 250 255

Glu Ala Ala Gln Arg Leu Gln His Arg Leu Ile Glu Pro Asn Thr Ala 260 265 270

Val Thr Arg Arg Phe Gly Ile Pro Gly Leu Lys Lys Thr Met Asp Trp 275 280 285

Phe Gly Tyr Tyr Gly Gly Pro Cys Arg Ala Pro Leu Gln Glu Leu Ser 290 295 300

Pro Ser Glu Glu Glu Ala Leu Arg Leu Asp Phe Ser Asn Asn Gly Trp 305 310 315

Leu

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Gly Leu Glu Leu Pro Val Asp Ala Val Val Thr Leu Ser Gln His Pro 35 40 45

Asn Ile Ile Gly Leu Lys Asp Ser Gly Gly Asp Val Thr Arg Thr Gly 50 60

Leu Ile Val His Lys Thr Ser Lys Gln Asp Phe Gln Val Leu Ala Gly 70 75 80

Ser Val Gly Phe Leu Leu Ala Ser Tyr Ala Val Gly Ala Val Gly Gly 85 90 95

Ile Val Gly Leu Ala Asn Val Leu Gly Ala Gln Val Cys Gln Leu Glu 100 105 110

Arg Leu Cys Leu Thr Gly Gln Gly Glu Ala Ala Gln Arg Leu Gln His 115 120 125

Arg Leu Ile Glu Pro Asn Thr Ala Val Thr Arg Arg Phe Gly Ile Pro 130 135 140

Gly Leu Lys Lys Thr Met Asp Trp Phe Gly Tyr Tyr Gly Gly Pro Cys 145 150 155 160

Arg Ala Pro Leu Xaa Glu Leu Ser Pro Ser Glu Glu Glu Ala Leu Arg 165 170 175

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Glu Leu Ser Gln Thr Leu Val Pro Thr Val 195 200

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<212> DNA

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eggaaceeg agaea;	geeca gaee	
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